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Mark Up Copy of Amendments to Claims

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Claim 74 has been amended as follows:

- 74. (Amended) A method for sequencing DNA which comprises:
 - treating the DNA with a mixture comprising an (a) oligonucleotide primer, a DNA polymerase, four different deoxynucleotides, and four different labeled dideoxynucleotides, under conditions permitting a deoxynucleotide or a labeled dideoxynucleotide or both to be incorporated into a DNA sequencing fragment, wherein each different deoxynucleotide and each different labeled dideoxynucleotide is complementary to one of the four nucleotides present in the wherein each labeled dideoxynucleotide comprises a chemical moiety attached via a linker to the dideoxynucleotide; and wherein each of the four different labeled dideoxynucleotides has a molecular weight which can be distinguished from the molecular weight ο£ the other three dideoxynucleotides using mass spectrometry;
 - (b) generating a plurality of DNA sequencing fragments having different lengths that are terminated with the labeled dideoxynucleotides so as to generate a plurality of different labeled DNA sequencing fragments, wherein each DNA sequencing fragment has a 3' end and the chemical moiety is attached via the linker to the 3' end of the DNA sequencing fragment;
 - (c) contacting the labeled DNA sequencing fragments with a surface coated with a

compound that specifically interacts with the chemical moiety attached via the linker to the end of the DNA sequencing fragments, thereby capturing the labeled DNA sequencing the surface[;], wherein the fragments on contacting is performed in a system comprising (i) a channel whose surface is coated with a compound that specifically interacts with the chemical moiety, wherein the channel comprises a plurality of ends, (ii) a plurality of wells each suitable for holding a sample, (iii) a connection between each end of the channel and a well, and (iv) a means for moving the sample through the channel between wells;

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- (d) washing the surface to remove non-bound components;
- (e) treating the labeled DNA sequencing fragments so as to release the labeled DNA sequencing fragments from the surface; and
- (f) determining the difference in molecular weight between different labeled DNA sequencing fragments which are represented as adjacent peaks on a mass spectra of the labeled DNA sequencing fragments produced using mass spectrometry, so as to sequence the DNA;

wherein either (i) the labeled dideoxynucleotides are biotinylated dideoxynucleotides selected from the group consisting of

where ddNTP1, ddNTP2, ddNTP3, and ddNTP4 represent four different dideoxynucleotides; or

(ii) the linker is selected from the group consisting of

CH₂NHC(0)CF₃

and

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